## Statistics Advanced - 2| Assignment

**Question 1:** What is hypothesis testing in statistics?

**Answer:** Hypothesis testing is a formal statistical procedure for deciding whether there is enough evidence in a sample of data to support a particular claim (hypothesis) about a population parameter. It starts with a null hypothesis ( $H_0$ ), defines an alternative hypothesis ( $H_1$  or  $H_2$ ), selects a test statistic and significance level ( $\alpha$ ), computes the statistic from the sample, and uses its sampling distribution to produce a p-value or compare to critical values. Based on that, we either reject  $H_0$  or fail to reject  $H_0$ , drawing conclusions with a controlled Type I error probability.

**Question 2:** What is the null hypothesis, and how does it differ from the alternative hypothesis?

**Answer:** Null hypothesis (H<sub>0</sub>): The default position — typically a statement of no effect or no difference (e.g.,  $\mu = \mu_0$ ).

Alternative hypothesis (H<sub>1</sub> or H<sub>a</sub>): The statement you want evidence for (e.g.,  $\mu \neq \mu_0$ ,  $\mu > \mu_0$ , or  $\mu < \mu_0$ ).

They are complementary. Hypothesis testing evaluates whether observed data are sufficiently incompatible with  $H_0$  to support  $H_1$ , at the chosen significance level.

**Question 3:** Explain the significance level in hypothesis testing and its role in deciding the outcome of a test.

**Answer:** The significance level  $\alpha$  is the threshold probability of making a Type I error (rejecting H<sub>0</sub> when it is actually true). Common  $\alpha$  values are 0.05 or 0.01. If the p-value  $\leq \alpha$ , we reject H<sub>0</sub>; otherwise, we fail to reject H<sub>0</sub>.  $\alpha$  therefore controls how strict we are about claiming a statistically significant result.

**Question 4:** What are Type I and Type II errors? Give examples of each.

**Answer:** Type I error ( $\alpha$ ): Rejecting H<sub>0</sub> when H<sub>0</sub> is true. Example: concluding a new drug works when it actually does not.

Type II error ( $\beta$ ): Failing to reject  $H_0$  when  $H_1$  is true. Example: concluding the new drug does not work when it actually does.

Power =  $1 - \beta$  is the probability of correctly rejecting a false  $H_0$ .

**Question 5:** What is the difference between a Z-test and a T-test? Explain when to use each.

**Answer:** Z-test: Used when the sampling distribution of the test statistic is normal and the population standard deviation ( $\sigma$ ) is known — or when the sample size is large and we approximate  $\sigma$  with the sample standard deviation. The test statistic uses the **normal** 

distribution.

plt.ylabel("Frequency")

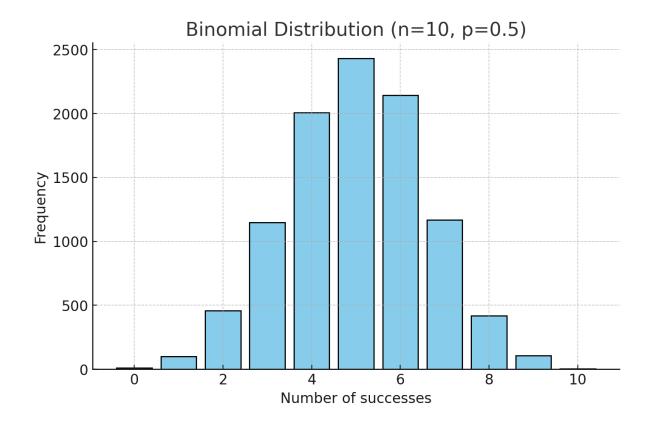
plt.grid(True) plt.show()

T-test: Used when  $\sigma$  is unknown and the sample comes from a normally distributed population (especially for small samples). The test statistic follows Student's t-distribution with n=1 degrees of freedom.

In practice: use a t-test when  $\sigma$  is unknown (most real situations); use a Z-test if  $\sigma$  is known or for very large samples where normal approximation is valid.

**Question 6:** Write a Python program to generate a binomial distribution with n=10 and p=0.5, then plot its histogram.

```
Answer:
import numpy as np
import matplotlib.pyplot as plt
# Parameters
n = 10
        # number of trials
p = 0.5
          # probability of success
trials = 10000 # number of experiments
# Generate binomial distribution
binom_samples = np.random.binomial(n=n, p=p, size=trials)
# Print summary statistics
print("Mean of samples:", binom_samples.mean())
print("Standard deviation of samples:", binom_samples.std(ddof=1))
# Plot histogram
plt.figure(figsize=(8,5))
plt.hist(binom_samples, bins=range(n+2), align='left', rwidth=0.8, color="skyblue",
edgecolor="black")
plt.title("Binomial Distribution (n=10, p=0.5)")
plt.xlabel("Number of successes")
```



**Question 7:** Implement hypothesis testing using *Z*-statistics for a sample dataset in Python. Show the Python code and interpret the results. sample\_data = [49.1, 50.2, 51.0, 48.7, 50.5, 49.8, 50.3, 50.7, 50.2, 49.6, 50.1, 49.9, 50.8, 50.4, 48.9, 50.6, 50.0, 49.7, 50.2, 49.5, 50.1, 50.3, 50.4, 50.5, 50.0, 50.7, 49.3, 49.8, 50.2, 50.9, 50.3, 50.4, 50.0, 49.7, 50.5, 49.9]

Answer: import numpy as np

from math import sqrt from scipy import stats import matplotlib.pyplot as plt

## # Sample data

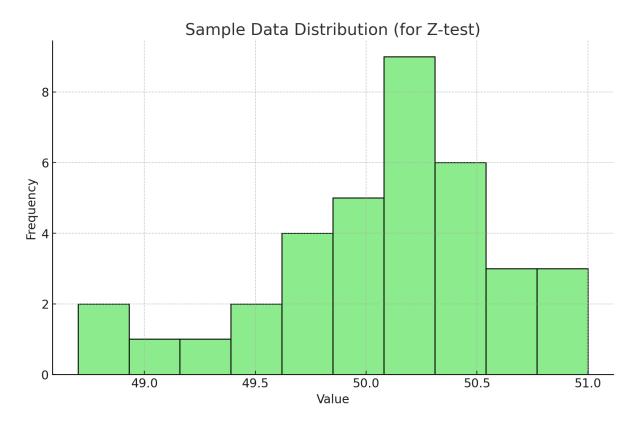
```
sample_data = [49.1, 50.2, 51.0, 48.7, 50.5, 49.8, 50.3, 50.7, 50.2, 49.6, 50.1, 49.9, 50.8, 50.4, 48.9, 50.6, 50.0, 49.7, 50.2, 49.5, 50.1, 50.3, 50.4, 50.5, 50.0, 50.7, 49.3, 49.8, 50.2, 50.9, 50.3, 50.4, 50.0, 49.7, 50.5, 49.9]
```

```
x = np.array(sample_data)
n = len(x)
sample_mean = x.mean()
sample_std = x.std(ddof=1)
```

## # Hypothesized population mean mu0 = 50

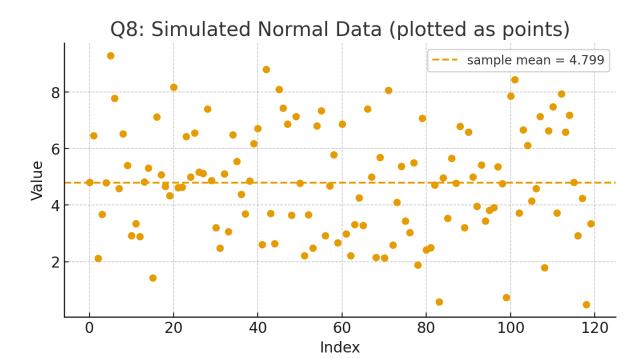
```
# Assume population sigma is known = 1.0 (for Z-test) sigma = 1.0
```

```
# Z-test statistic
z_stat = (sample_mean - mu0) / (sigma / sqrt(n))
# Two-tailed p-value
p val = 2 * (1 - stats.norm.cdf(abs(z stat)))
print("Sample size:", n)
print("Sample mean:", sample_mean)
print("Sample std dev:", sample_std)
print("Z-statistic:", z_stat)
print("p-value:", p_val)
# Interpretation
alpha = 0.05
if p val < alpha:
  print("Reject H0: Evidence suggests the mean is different from 50.")
else:
  print("Fail to reject H0: No strong evidence mean differs from 50.")
# Plot histogram of sample data
plt.hist(x, bins=10, color="lightgreen", edgecolor="black")
plt.title("Sample Data Distribution (for Z-test)")
plt.xlabel("Value")
plt.ylabel("Frequency")
plt.grid(True)
plt.show()
```



**Question 8:** Write a Python script to simulate data from a normal distribution and calculate the 95% confidence interval for its mean. Plot the data using Matplotlib.

```
Answer: import numpy as np
import matplotlib.pyplot as plt
from scipy import stats
from math import sqrt
# Parameters for the normal distribution
np.random.seed(42)
                        # for reproducibility
mu_true = 5.0
                     # true mean
                      # true standard deviation
sigma true = 2.0
n = 120
                  # sample size
# Simulate data
data = np.random.normal(loc=mu_true, scale=sigma_true, size=n)
# Sample statistics
sample_mean = data.mean()
sample_std = data.std(ddof=1)
se = sample std / sqrt(n)
# 95% confidence interval using t-distribution
alpha = 0.05
t crit = stats.t.ppf(1 - alpha/2, df=n-1)
ci_lower = sample_mean - t_crit * se
ci_upper = sample_mean + t_crit * se
print("Sample mean:", sample_mean)
print("Sample standard deviation:", sample std)
print("95% Confidence Interval: [{:.4f}, {:.4f}]".format(ci_lower, ci_upper))
# Plot the data
plt.figure(figsize=(8,5))
plt.plot(data, marker='o', linestyle='none', alpha=0.7, color="purple")
plt.axhline(sample_mean, color="red", linestyle="--", label=f"Sample Mean =
{sample_mean:.2f}")
plt.axhline(ci_lower, color="blue", linestyle="--", label=f"95% CI_Lower = {ci_lower:.2f}")
plt.axhline(ci_upper, color="blue", linestyle="--", label=f"95% CI Upper = {ci_upper:.2f}")
plt.title("Simulated Normal Data with 95% Confidence Interval")
plt.xlabel("Observation Index")
plt.ylabel("Value")
plt.legend()
plt.grid(True)
plt.show()
```



**Question 9:** Write a Python function to calculate the Z-scores from a dataset and visualize the standardized data using a histogram. Explain what the Z-scores represent in terms of standard deviations from the mean.

Answer: import numpy as np import matplotlib.pyplot as plt # Function to calculate Z-scores def calculate\_z\_scores(data): arr = np.array(data)mean = arr.mean() std = arr.std(ddof=0) # population standard deviation z scores = (arr - mean) / std return z\_scores # Example dataset (you can replace with any data) sample\_data = [49.1, 50.2, 51.0, 48.7, 50.5, 49.8, 50.3, 50.7, 50.2, 49.6, 50.1, 49.9, 50.8, 50.4, 48.9, 50.6, 50.0, 49.7, 50.2, 49.5, 50.1, 50.3, 50.4, 50.5, 50.0, 50.7, 49.3, 49.8, 50.2, 50.9, 50.3, 50.4, 50.0, 49.7, 50.5, 49.9] # Calculate Z-scores zs = calculate\_z\_scores(sample\_data) # Print first few Z-scores print("First 10 Z-scores:", np.round(zs[:10], 3)) print("Mean of Z-scores:", np.round(zs.mean(), 5)) print("Std of Z-scores:", np.round(zs.std(ddof=0), 5))

```
# Plot histogram
plt.figure(figsize=(8,5))
plt.hist(zs, bins=10, color="orange", edgecolor="black")
plt.title("Histogram of Z-scores (Standardized Data)")
plt.xlabel("Z-score")
plt.ylabel("Frequency")
plt.grid(True)
plt.show()
```

